

Notice of References Cited	Application/Control No. 10/771,417	Applicant(s)/Patent Under Reexamination WATANABE ET AL.	
	Examiner Bridget E. Bunner	Art Unit 1647	Page 1 of 3

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	L	US-			
	M	US-			

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	V	Bork, A. Powers and pitfalls in sequence analysis: the 70% hurdle. Genome Res 10: 398-400, 2000.
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	X	Smith et al. The challenges of genome sequence annotation or "The devil is in the details". Nature Biotech 15: 1222-1223, 1997.

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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	X	Ngo et al. Computational complexity, protein structure prediction, and the Levinthal paradox. The Protein Folding Problem and Tertiary Structure Prediction, pp. 492-495, 1994.

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Examiner

Bridget E. Bunner

2/28/06
H/b/b

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